OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/897,844A

DATE: 11/23/2001
TIME: 10:57:17

Input Set: N:\Crf3\RULE60\09897844A.RAW
Output Set: N:\CRF3\11232001\1897844A.raw

```
1 <110> APPLICANT: Cox III, George Norbert
                                                                  ENTERED
            Case, Casey Christopher
            Eisenberg, Stephen P.
    3
            Jarvis, Eric Edward
    4
            Spratt, Sharon Kaye
            Sangamo Biosciences, Inc.
    6
    7 <120> TITLE OF INVENTION: Regulation of Endogenous Gene Expression in Cells
            Using Zinc Finger Proteins
    9 <130> FILE REFERENCE: 019496-002200US
    10 <140> CURRENT APPLICATION NUMBER: 09/897,844A
    11 <141> CURRENT FILING DATE: 2001-07-02
    12 <150> PRIOR APPLICATION NUMBER: US/09/229,037
    13 <151> PRIOR FILING DATE: 1999-01-12
    14 <160> NUMBER OF SEQ ID NOS: 40
    15 <170> SOFTWARE: PatentIn Ver. 2.0
    17 <210> SEQ ID NO: 1
    18 <211> LENGTH: 25
    19 <212> TYPE: PRT
    20 <213> ORGANISM: Artificial Sequence
    21 <220> FEATURE:
    22 <223> OTHER INFORMATION: Description of Artificial Sequence: exemplary motif
            of C2H2 class of zinc finger proteins (ZFP)
    24 <220> FEATURE:
    25 <221> NAME/KEY: MOD_RES
    26 <222> LOCATION: (2)..(3)
    27 <223> OTHER INFORMATION: Xaa = any amino acid
    28 <220> FEATURE:
    29 <221> NAME/KEY: MOD_RES
    30 <222> LOCATION: (4)..(5)
    31 <223> OTHER INFORMATION: Xaa = any amino acid, may be present or absent
    32 <220> FEATURE:
    33 <221> NAME/KEY: MOD_RES
    34 <222> LOCATION: (7)..(18)
    35 <223> OTHER INFORMATION: Xaa = any amino acid
    36 <220> FEATURE:
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    38 <222> LOCATION: (20)..(22)
    39 <223> OTHER INFORMATION: Xaa = any amino acid
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    41 <221> NAME/KEY: MOD_RES
    42 <222> LOCATION: (23)..(24)
    43 <223> OTHER INFORMATION: Xaa = any amino acid, may be present or absent
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    46
             Xaa Xaa His Xaa Xaa Xaa Xaa His
W--> 47
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Input Set: N:\Crf3\RULE60\09897844A.RAW
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   52 <212> TYPE: DNA
   53 <213> ORGANISM: Artificial Sequence
   55 <223> OTHER INFORMATION: Description of Artificial Sequence: ZFP target site
            with two overlapping D-able subsites
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   57 <220> FEATURE:
   58 <221> NAME/KEY: modified_base
   59 <222> LOCATION: (1)..(2)
   60 <223> OTHER INFORMATION: n = g,a,c or t
    61 <220> FEATURE:
    62 <221> NAME/KEY: modified_base
    63 <222> LOCATION: (5)
    64 <223> OTHER INFORMATION: n = g,a,c or t
    65 <220> FEATURE:
    66 <221> NAME/KEY: modified_base
    67 <222> LOCATION: (8)
    68 <223> OTHER INFORMATION: n = g,a,c or t
    69 <220> FEATURE:
    70 <221> NAME/KEY: modified_base
    72 <223> OTHER INFORMATION: n = a,c or t; if g, then position 10 cannot be g
             or t
    73
    74 <220> FEATURE:
    75 <221> NAME/KEY: modified_base
    77 <223> OTHER INFORMATION: n = a or c; if g or t, then position 9 cannot be g
     78 <400> SEQUENCE: 2
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w--> 79
     81 <210> SEQ ID NO: 3
     82 <211> LENGTH: 10
     83 <212> TYPE: DNA
     84 <213> ORGANISM: Artificial Sequence
     86 <223> OTHER INFORMATION: Description of Artificial Sequence: ZFP target site
     85 <220> FEATURE:
              with three overlapping D-able subsites
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     88 <220> FEATURE:
     89 <221> NAME/KEY: modified_base
     90 <222> LOCATION: (1)..(2)
     91 <223> OTHER INFORMATION: n = g,a,c or t
     92 <220> FEATURE:
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      94 <222> LOCATION: (5)
     95 <223> OTHER INFORMATION: n = g,a,c or t
      96 <220> FEATURE:
      97 <221> NAME/KEY: modified_base
      98 <222> LOCATION: (8)
      99 <223> OTHER INFORMATION: n = g,a,c or t
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     104 <211> LENGTH: 5
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     106 <213> ORGANISM: Artificial Sequence
     108 <223> OTHER INFORMATION: Description of Artificial Sequence:linker
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     113 <210> SEQ ID NO: 5
     114 <211> LENGTH: 5
      115 <212> TYPE: PRT
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     118 <223> OTHER INFORMATION: Description of Artificial Sequence: linker
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                Thr Gly Glu Lys Pro
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      123 <210> SEQ ID NO: 6
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      125 <212> TYPE: PRT
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       135 <212> TYPE: PRT
       136 <213> ORGANISM: Artificial Sequence
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       141
       143 <210> SEQ ID NO: 8
       144 <211> LENGTH: 5
       145 <212> TYPE: PRT
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       149 <400> SEQUENCE: 8
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       153 <210> SEQ ID NO: 9
       154 <211> LENGTH: 8
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156 <213> ORGANISM: Artificial Sequence
158 <223> OTHER INFORMATION: Description of Artificial Sequence:linker
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164 <211> LENGTH: 9
165 <212> TYPE: PRT
166 <213> ORGANISM: Artificial Sequence
168 <223> OTHER INFORMATION: Description of Artificial Sequence:linker
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170
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171
173 <210> SEQ ID NO: 11
174 <211> LENGTH: 12
175 <212> TYPE: PRT
 176 <213> ORGANISM: Artificial Sequence
 178 <223> OTHER INFORMATION: Description of Artificial Sequence:linker
 179 <400> SEQUENCE: 11
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 183 <210> SEQ ID NO: 12
 184 <211> LENGTH: 16
 185 <212> TYPE: PRT
 186 <213> ORGANISM: Artificial Sequence
 188 <223> OTHER INFORMATION: Description of Artificial Sequence:linker
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  189 <400> SEQUENCE: 12
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  193 <210> SEQ ID NO: 13
  194 <211> LENGTH: 25
  195 <212> TYPE: DNA
  196 <213> ORGANISM: Artificial Sequence
  198 <223> OTHER INFORMATION: Description of Artificial Sequence: ZFP target site
            region surrounding initiation site of vascular
            endothelial growth factor (VEGF) gene containing
  199
  200
            two 9-base pair target sites
  201
  202 <220> FEATURE:
  203 <221> NAME/KEY: protein_bind
  204 <222> LOCATION: (4)..(12)
  205 <223> OTHER INFORMATION: upstream 9-base pair ZFP VEGF1 target site
  206 <220> FEATURE:
  207 <221> NAME/KEY: protein_bind
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Input Set : N:\Crf3\RULE60\09897844A.RAW
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208 <222> LOCATION: (14)..(22) 209 <223> OTHER INFORMATION: downstream 9-base pair ZFP VEGF3a target site 210 <400> SEQUENCE: 13 25 agcggggagg atcgcggagg cttgg 213 <210> SEQ ID NO: 14 214 <211> LENGTH: 298 215 <212> TYPE: DNA 216 <213> ORGANISM: Artificial Sequence 218 <223> OTHER INFORMATION: Description of Artificial Sequence: VEGF1 ZFP construct targeting upstream 9-base pair target 219 site in VEGF promoter 220 221 <220> FEATURE: 222 <221> NAME/KEY: CDS 223 <222> LOCATION: (2)..(298) 224 <223> OTHER INFORMATION: VEGF1 g gta ccc ata cct ggc aag aag cag cac atc tgc cac atc cag ggc 49 225 <400> SEQUENCE: 14 Val Pro Ile Pro Gly Lys Lys Gln His Ile Cys His Ile Gln Gly 226 227 tgt ggt aaa gtt tac ggc aca acc tca aat ctg cgt cgt cac ctg cgc 97 228 Cys Gly Lys Val Tyr Gly Thr Thr Ser Asn Leu Arg Arg His Leu Arg 229 230 tgg cac acc ggc gag agg cct ttc atg tgt acc tgg tcc tac tgt ggt 20 145 231 Trp His Thr Gly Glu Arg Pro Phe Met Cys Thr Trp Ser Tyr Cys Gly 232 233 40 aaa cgc ttc acc cgt tcg tca aac ctg cag cgt cac aag cgt acc cac 193 234 Lys Arg Phe Thr Arg Ser Ser Asn Leu Gln Arg His Lys Arg Thr His 235 236 55 acc ggt gag aag aaa ttt gct tgc ccg gag tgt ccg aag cgc ttc atg 241 237 Thr Gly Glu Lys Lys Phe Ala Cys Pro Glu Cys Pro Lys Arg Phe Met 238 239 75 70 cgt agt gac cac ctg tcc cgt cac atc aag acc cac cag aat aag aag 289 240 Arg Ser Asp His Leu Ser Arg His Ile Lys Thr His Gln Asn Lys Lys 241 242 90 85 298 243 ggt gga tcc 244 Gly Gly Ser 245 247 <210> SEQ ID NO: 15 248 <211> LENGTH: 99 249 <212> TYPE: PRT 250 <213> ORGANISM: Artificial Sequence 252 <223> OTHER INFORMATION: Description of Artificial Sequence: VEGF1 ZFP 251 <220> FEATURE: construct targeting upstream 9-base pair target site in VEGF promoter 254 255 <400> SEQUENCE: 15 Val Pro Ile Pro Gly Lys Lys Gln His Ile Cys His Ile Gln Gly 256 10 Cys Gly Lys Val Tyr Gly Thr Thr Ser Asn Leu Arg Arg His Leu Arg 257 258

VERIFICATION SUMMARY

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Input Set : N:\Crf3\RULE60\09897844A.RAW
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L:45 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 L:47 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 L:101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3